

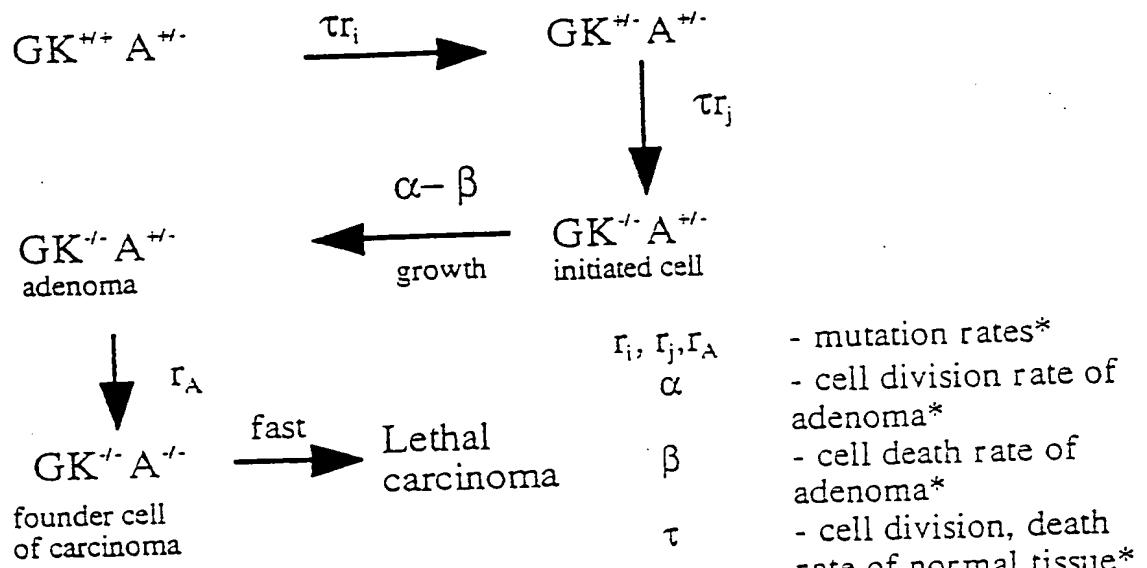


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Fig. 1



* all variables may be affected by exposure, xenometabolism, DNA repair



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Fig. 2A

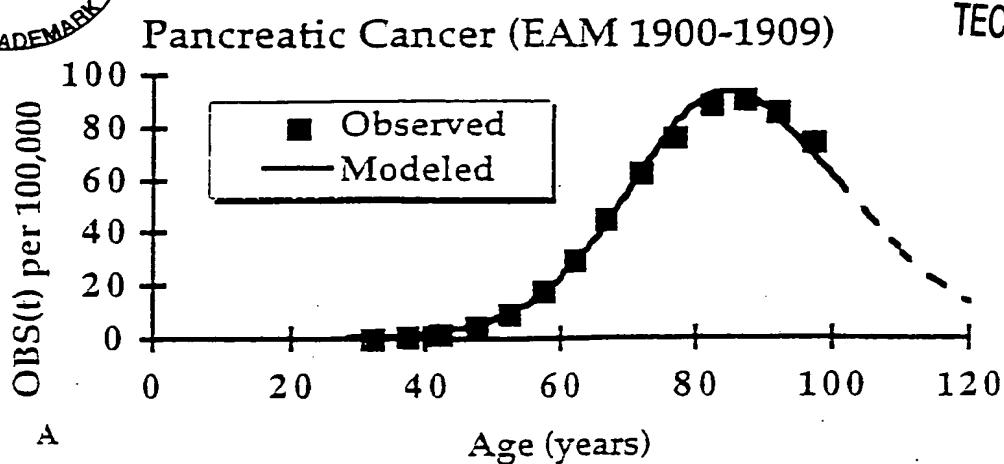
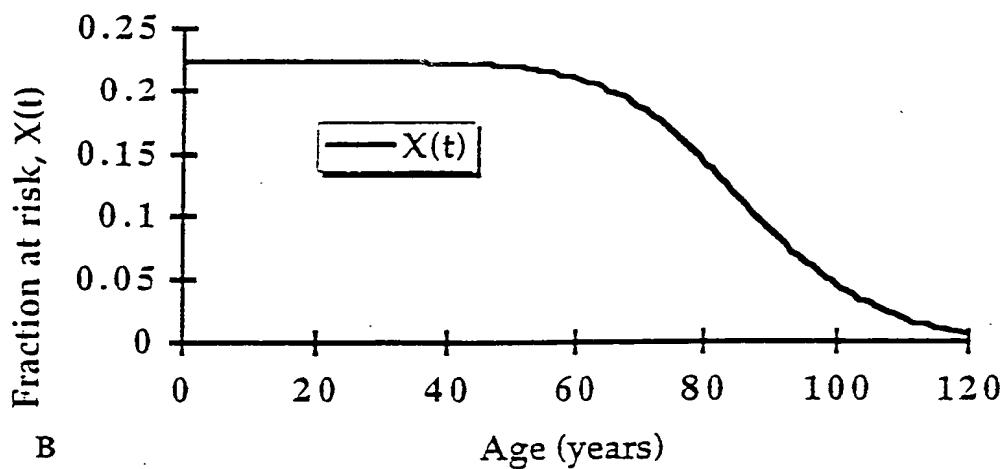


Fig. 2B



App No.: 09/503,758
Title: Methods of Identifying Point
Inventor: William G. Thilly

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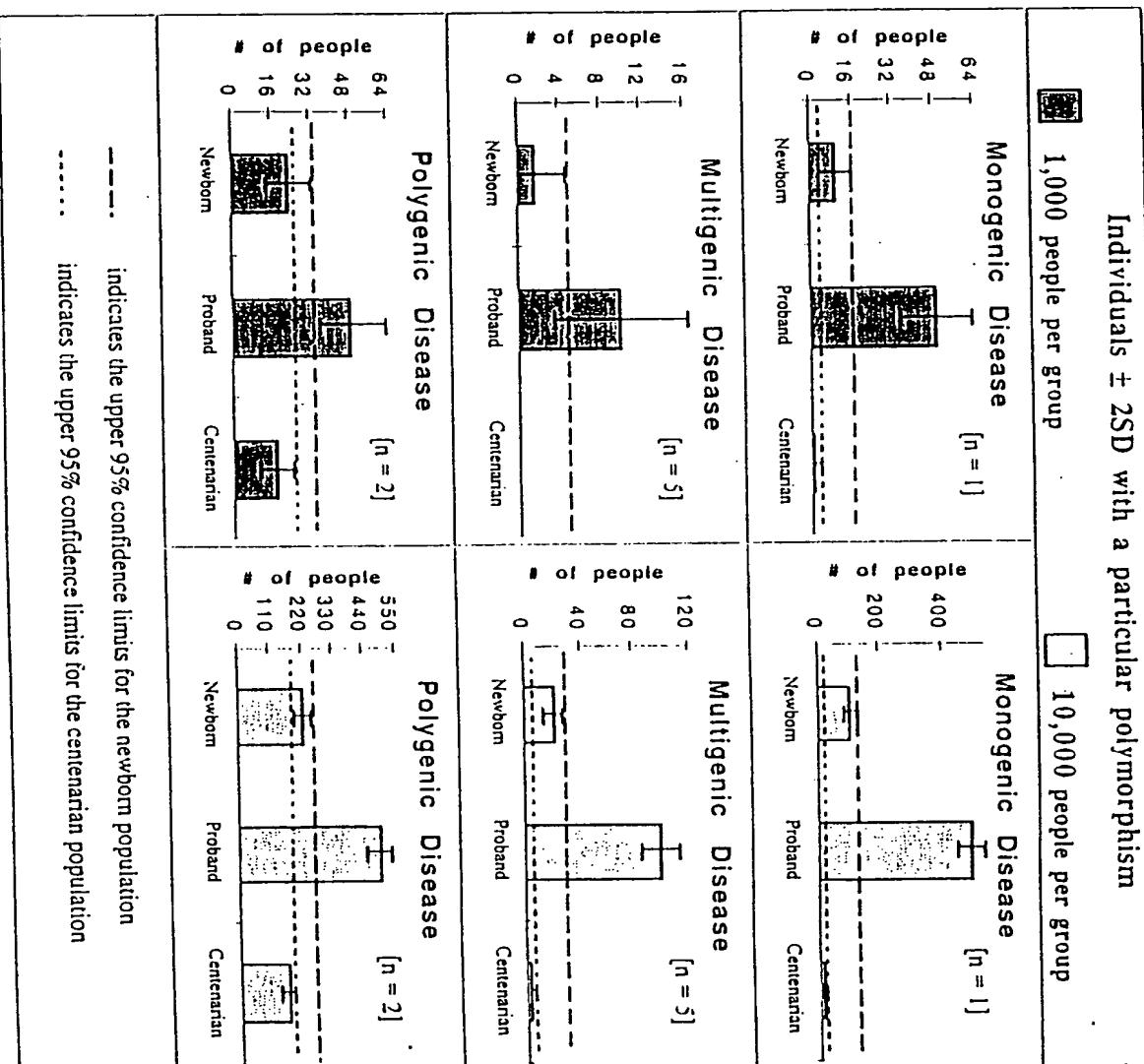
Fig. 30

Fig. 3A

Individuals \pm 2SD with a particular polymorphism

1,000 people per group

10,000 people per group



— — — — — indicates the upper 95% confidence limits for the newborn population
 - - - - - indicates the upper 95% confidence limits for the centenarian population

Fig. 3F

Fig. 3D



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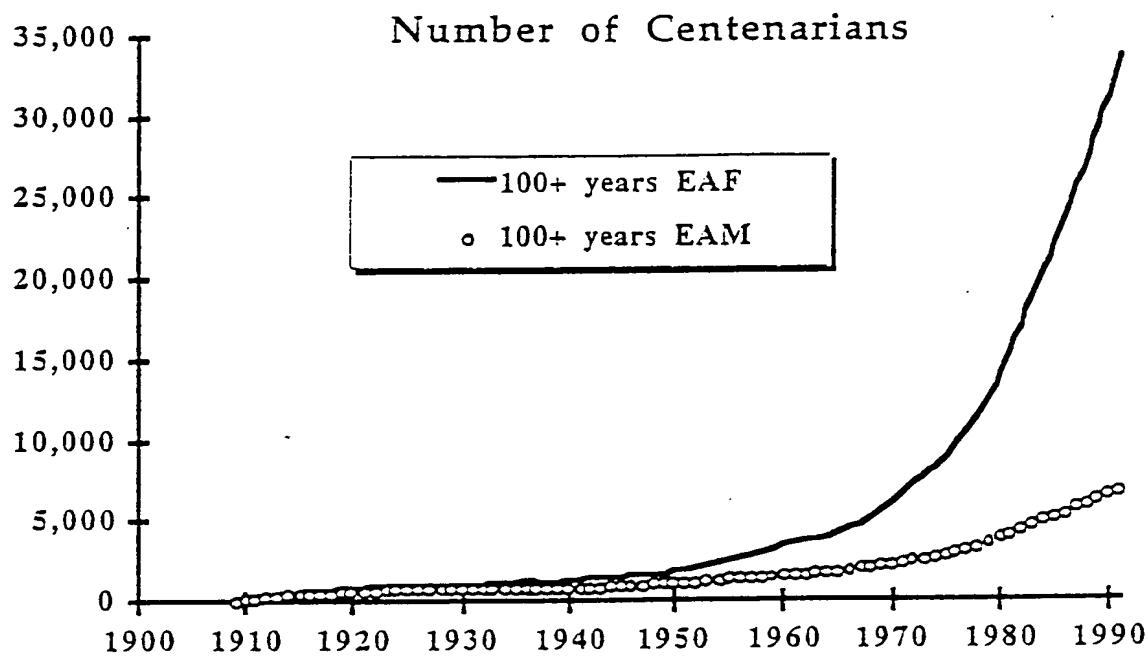


Fig. 4



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Fig. 5A

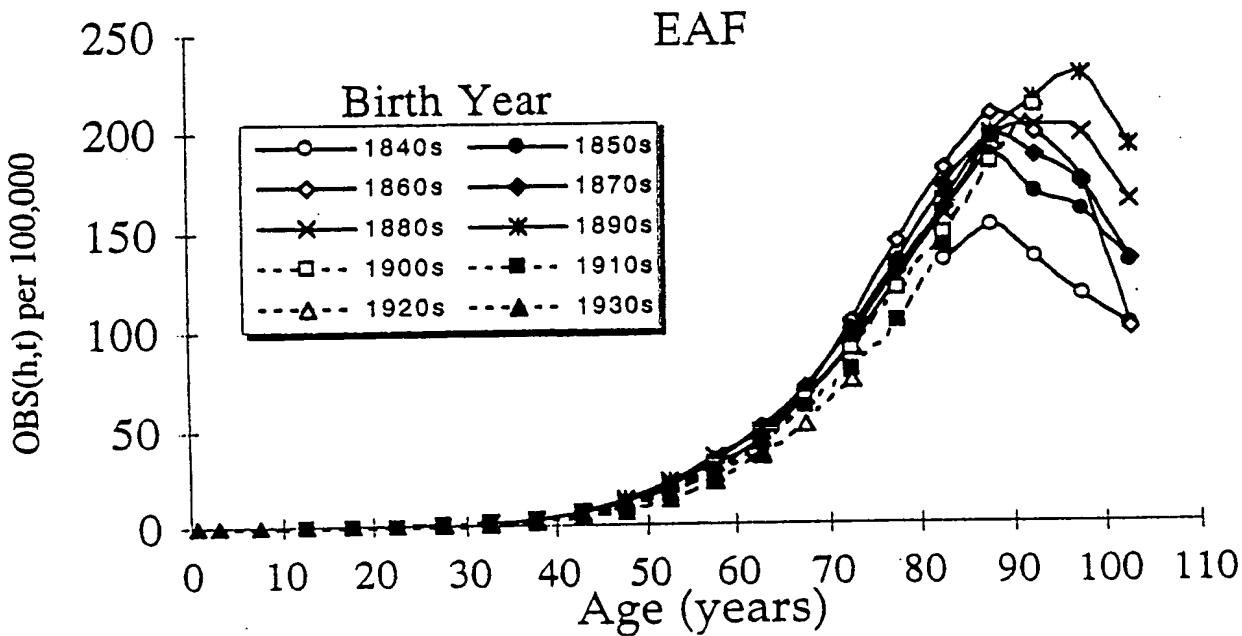
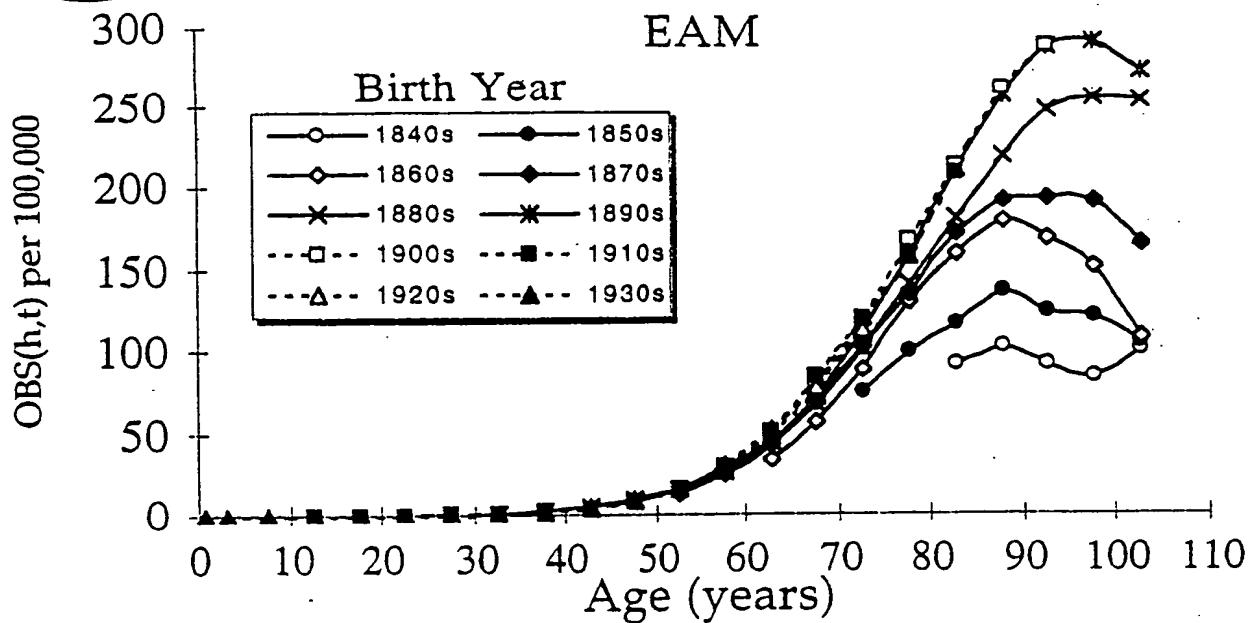


Fig. 5B

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Fig. 6A

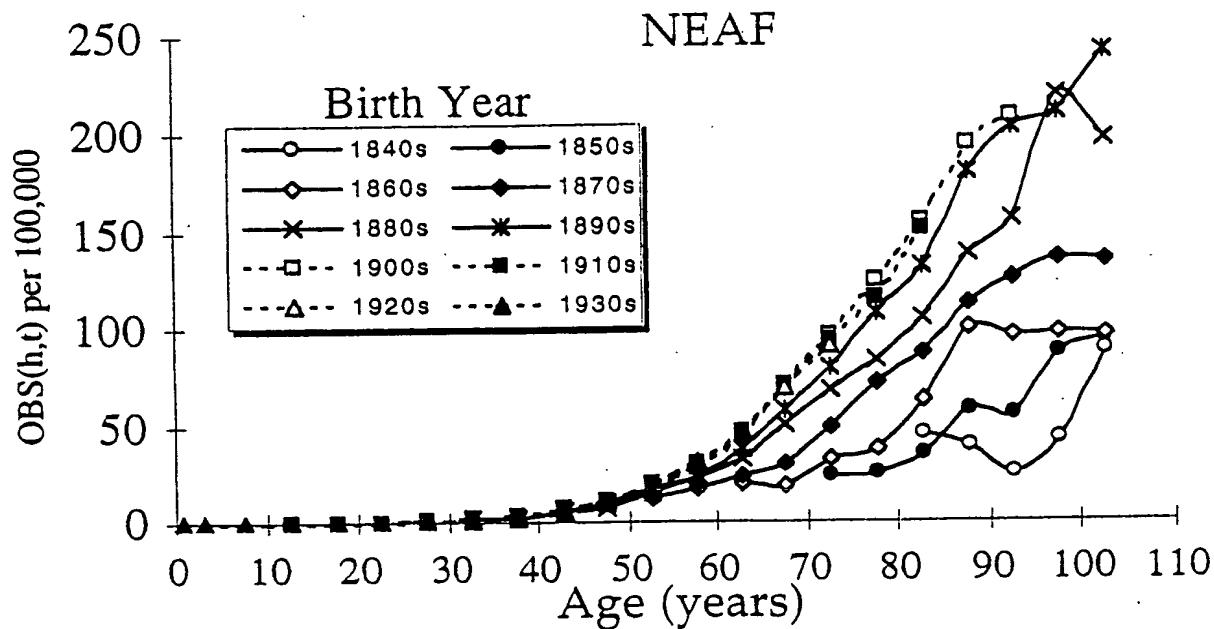
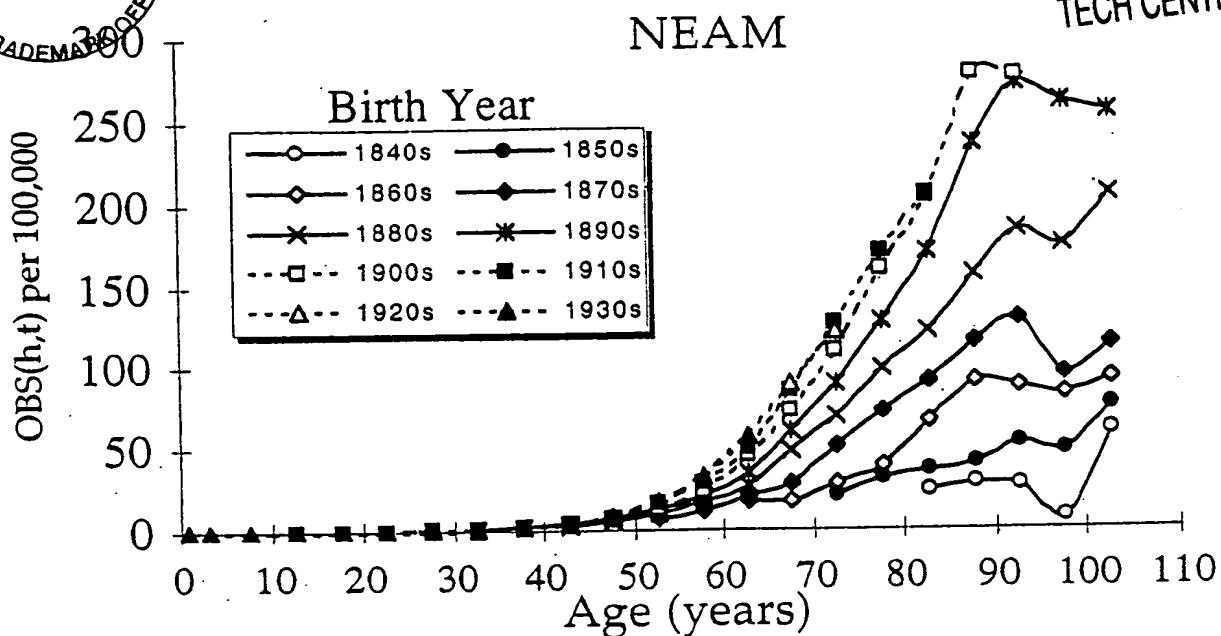


Fig. 6B



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Fig. 7A

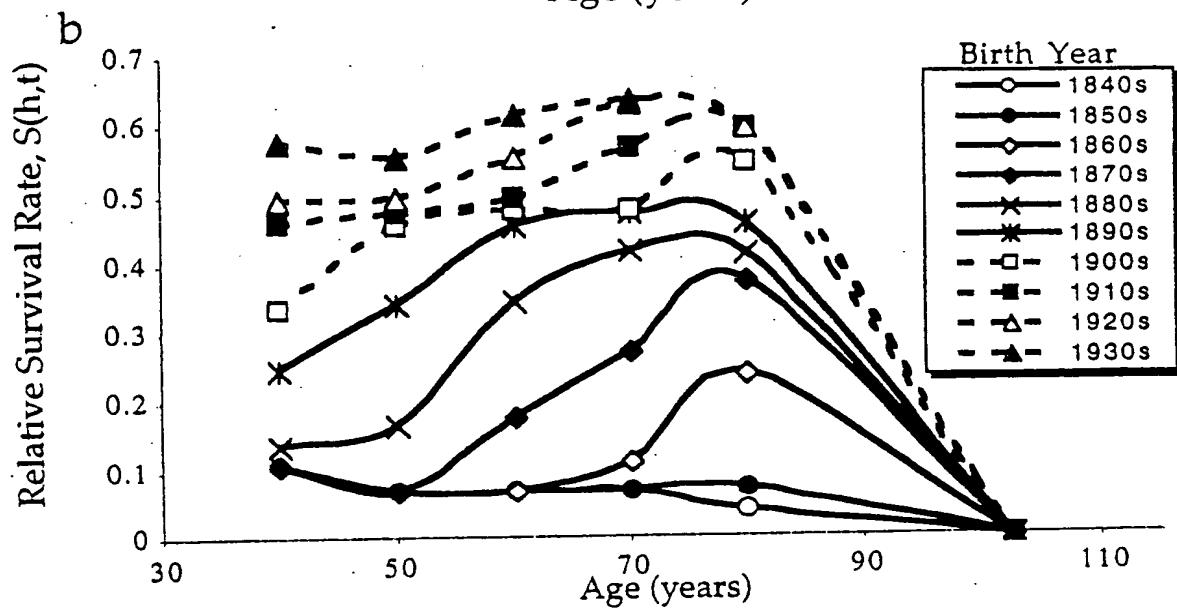
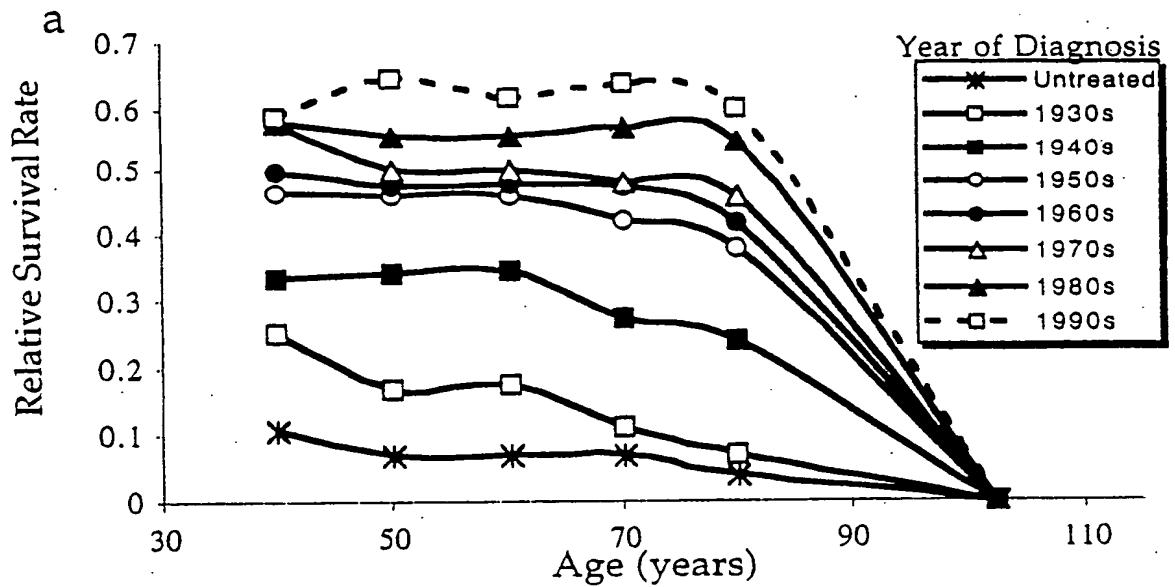


Fig. 7B

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Inventor: William G. Thilly

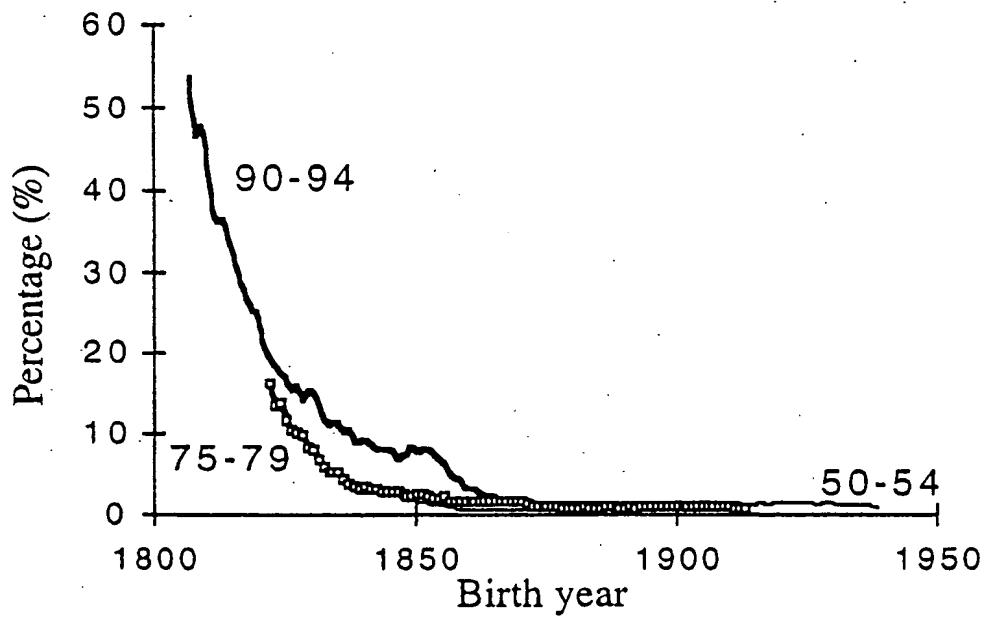
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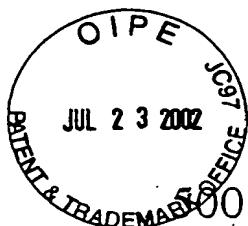
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Fig. 8





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Fig. 9A

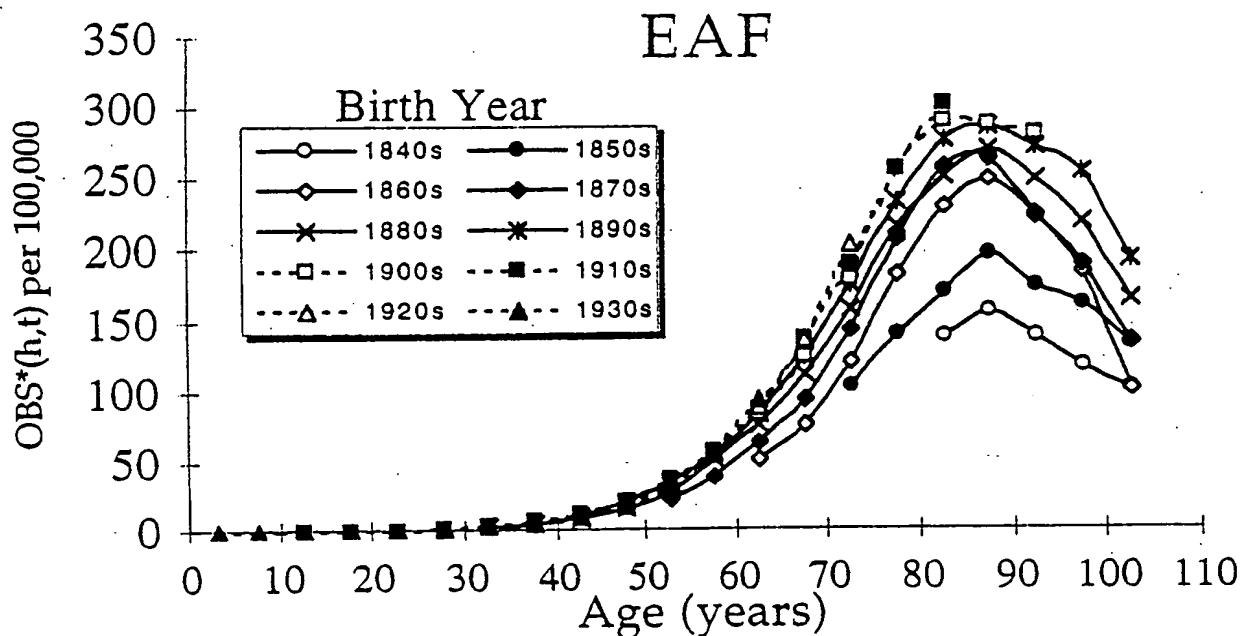
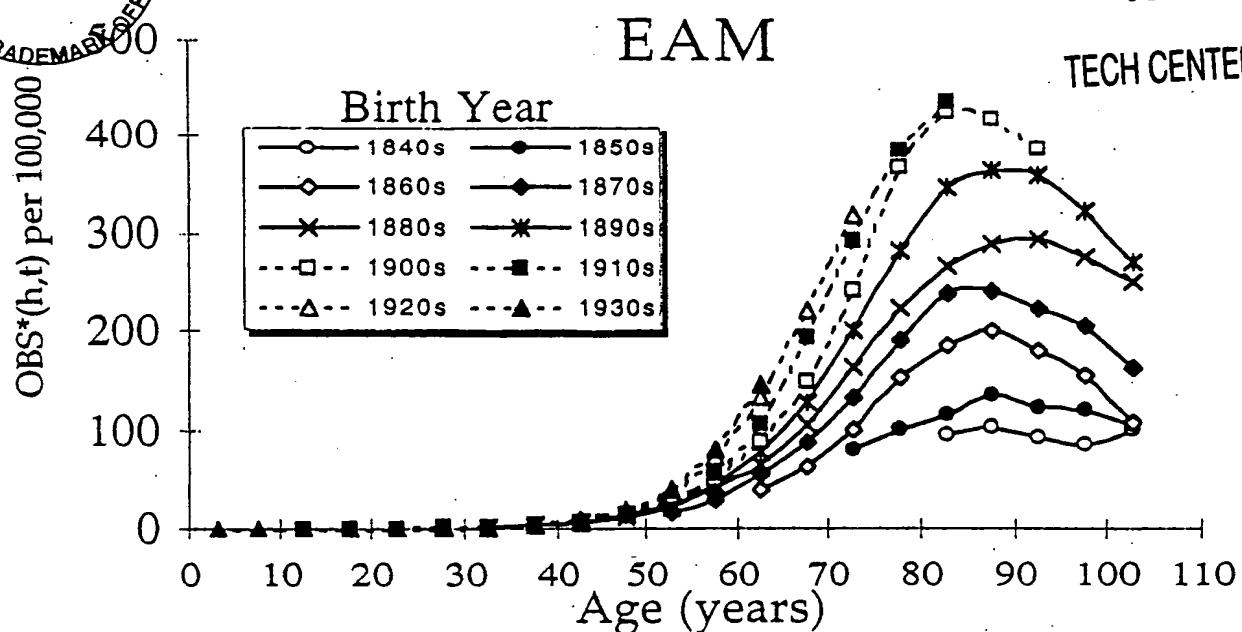
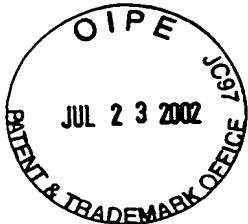


Fig. 9B

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Fig. 10A

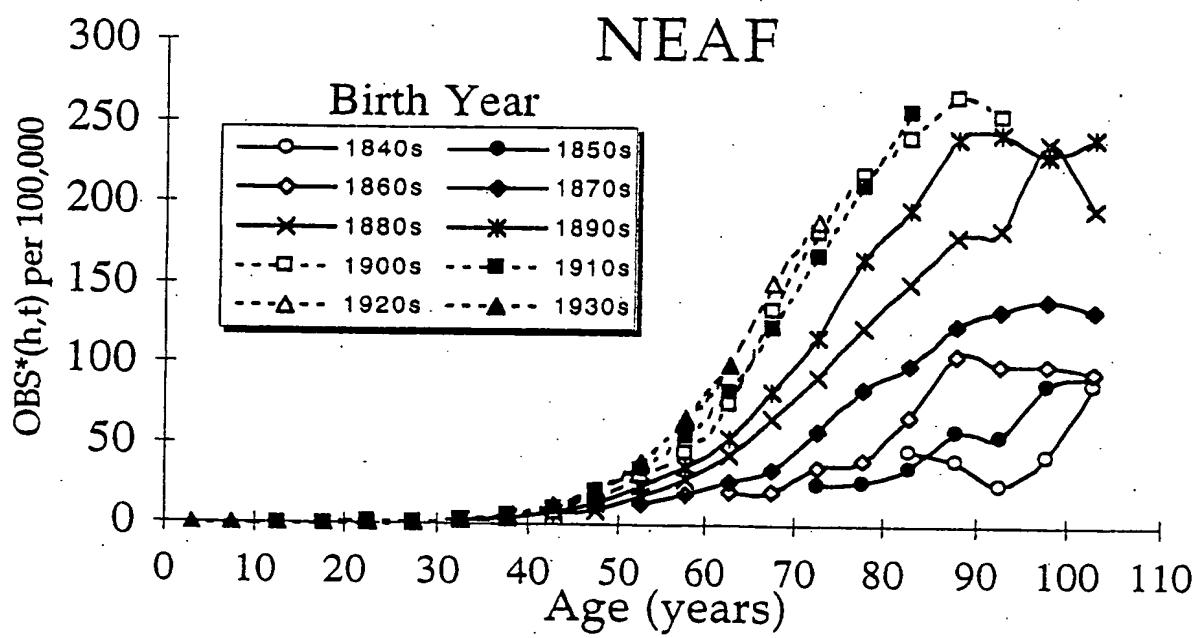
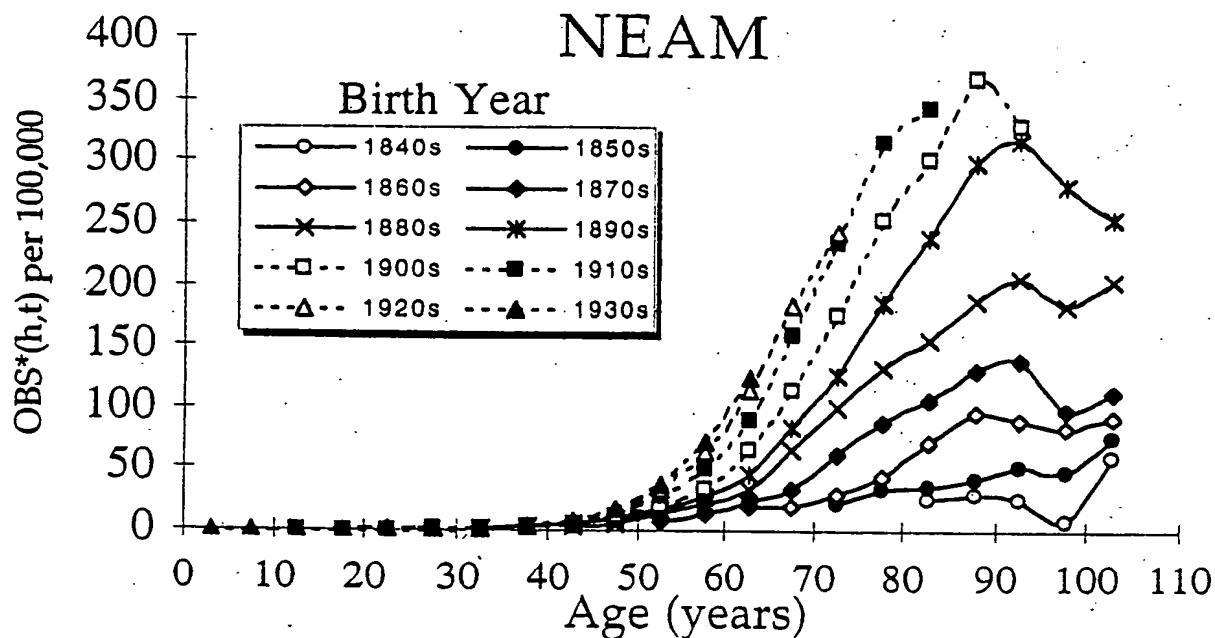


Fig. 10B

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Fig. 11A

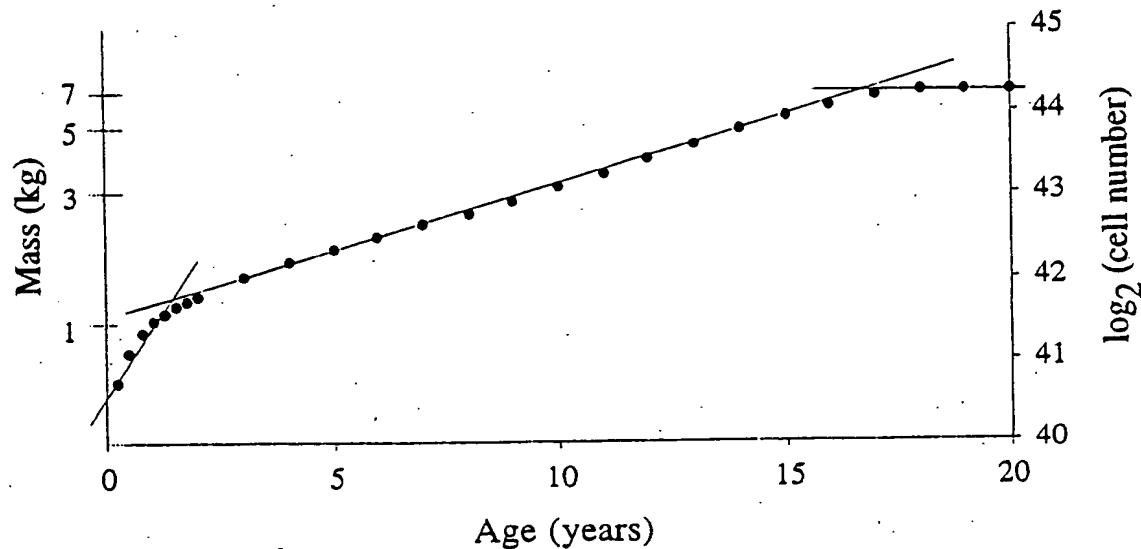
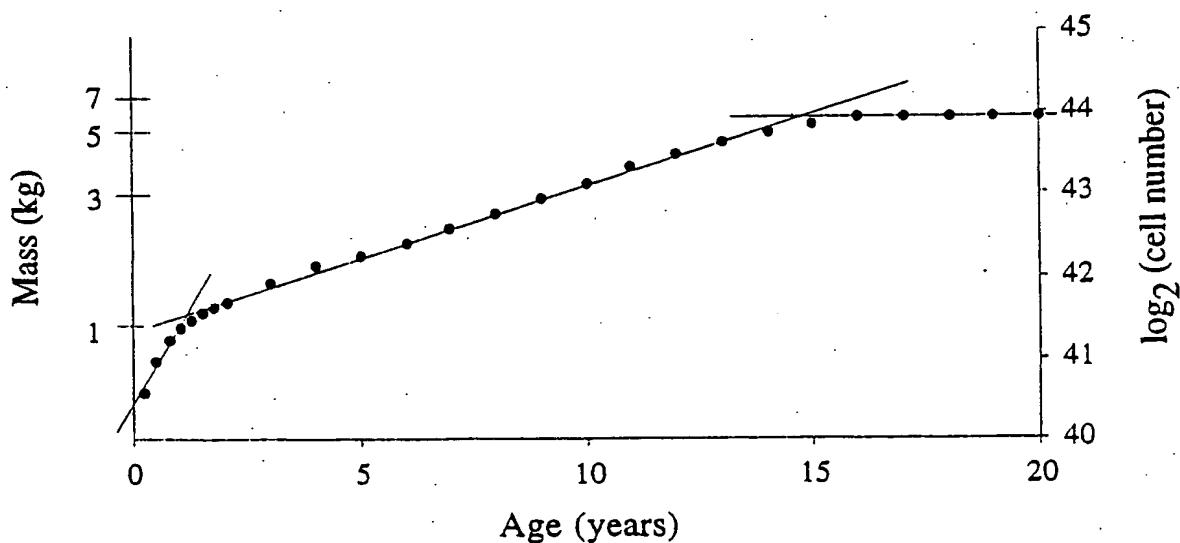


Fig. 11B



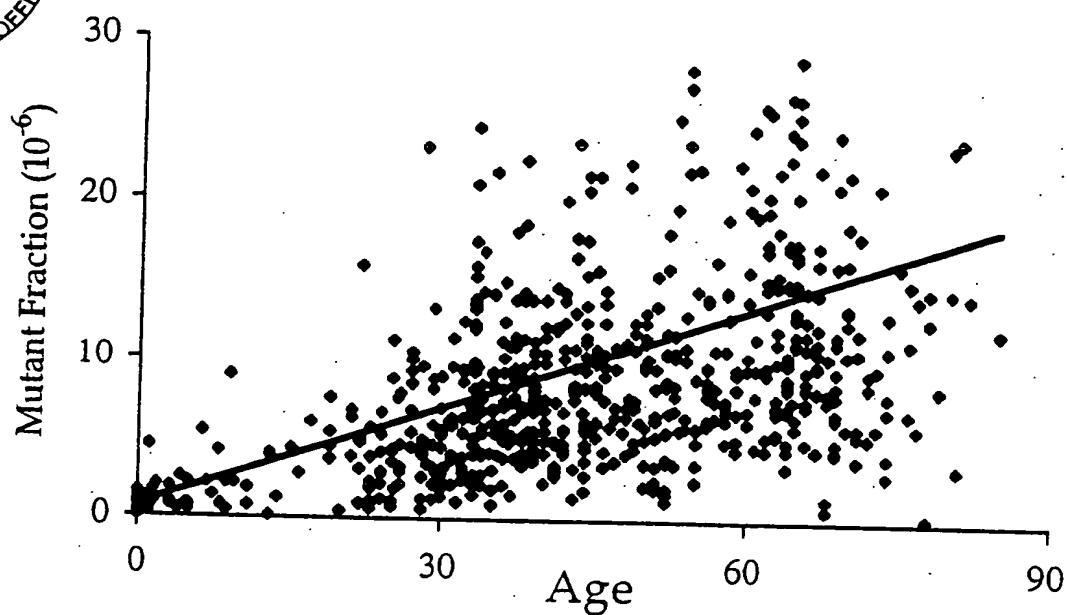


Fig. 12

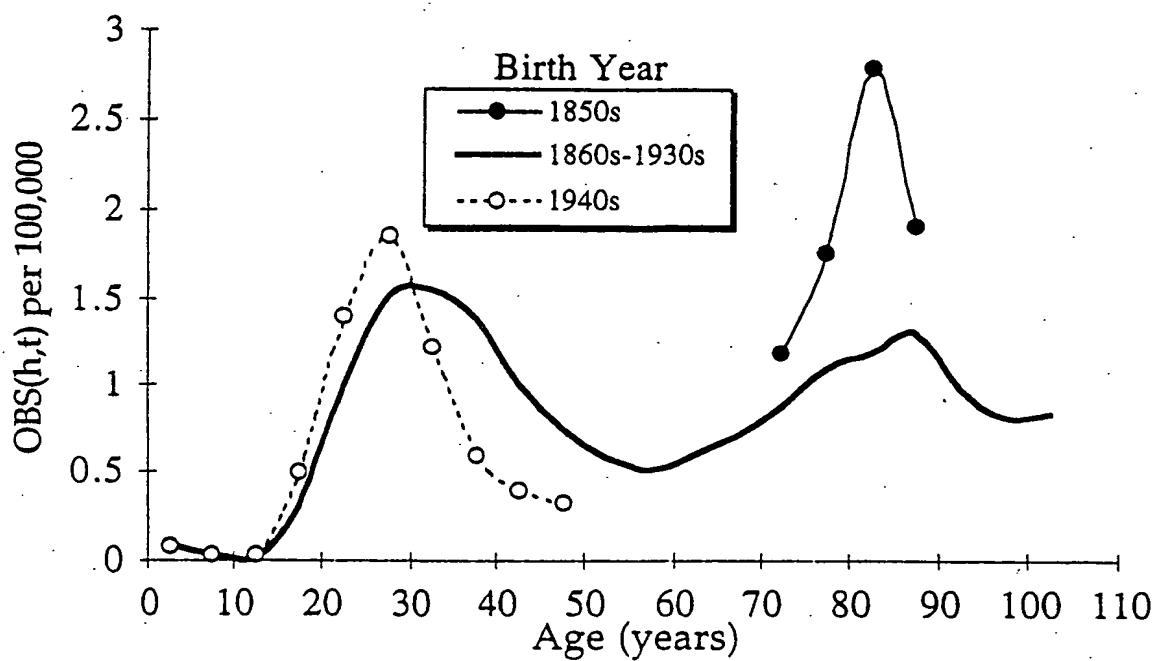
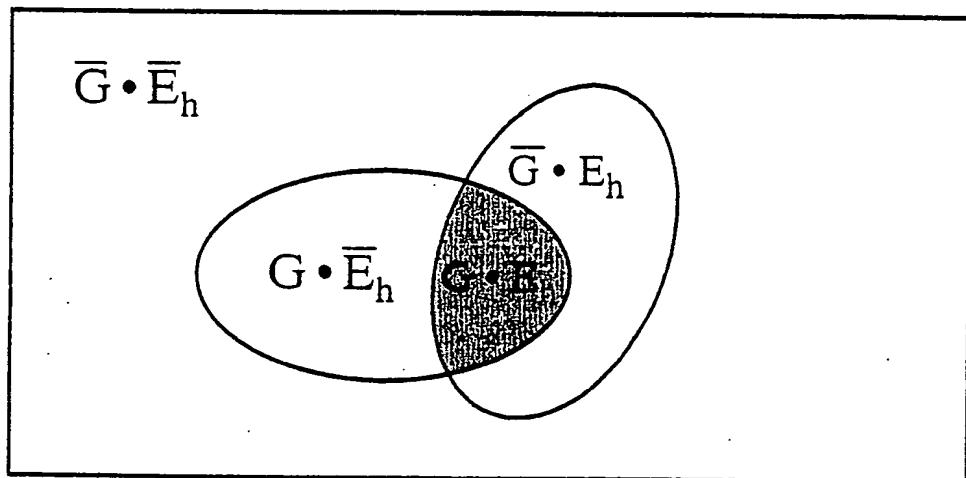


Fig. 13



Fig. 14



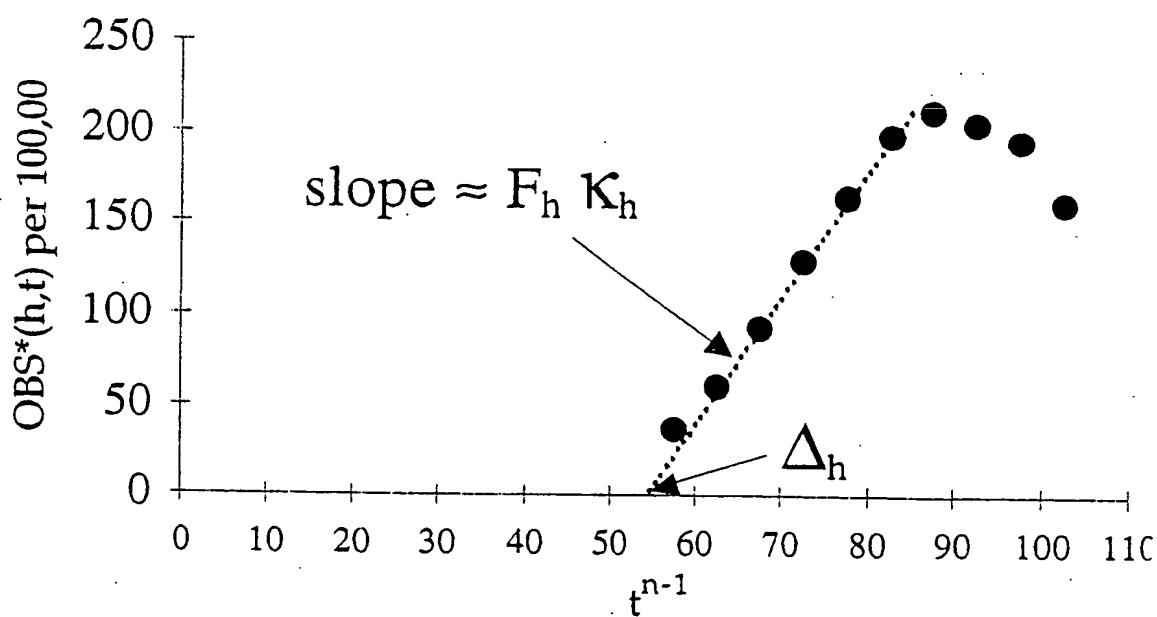
$G = F(h,t)_{\text{genetic}}$

$E_h = F(h,t)_{\text{environmental}}$

$\bar{G} = (1 - G) = \text{not in } F(h,t)_{\text{genetic}}$

$\bar{E}_h = (1 - E_h) = \text{not in } F(h,t)_{\text{environmental}}$

Fig. 15





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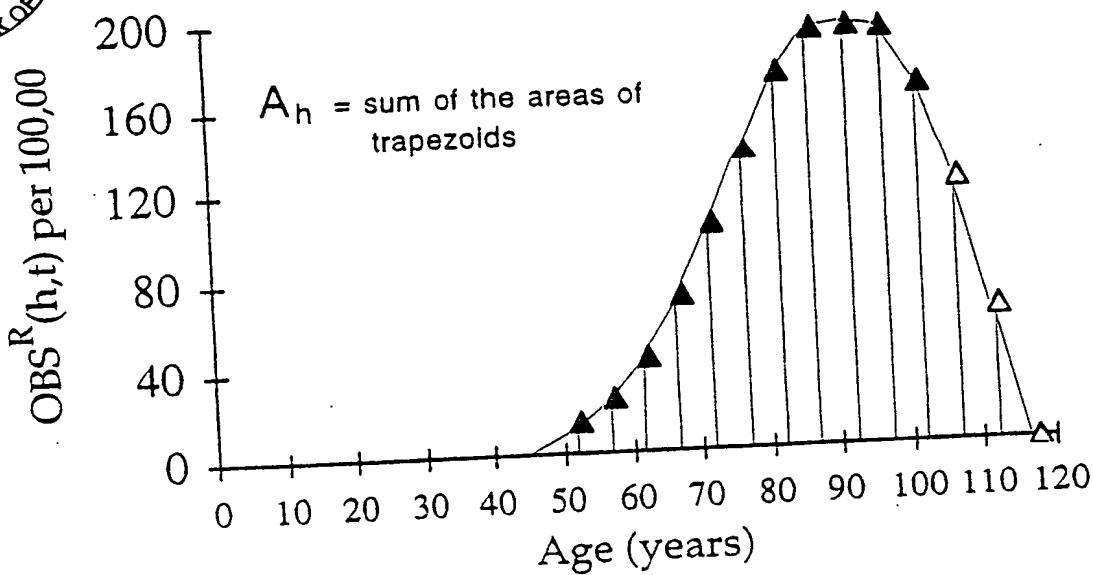


Fig. 16

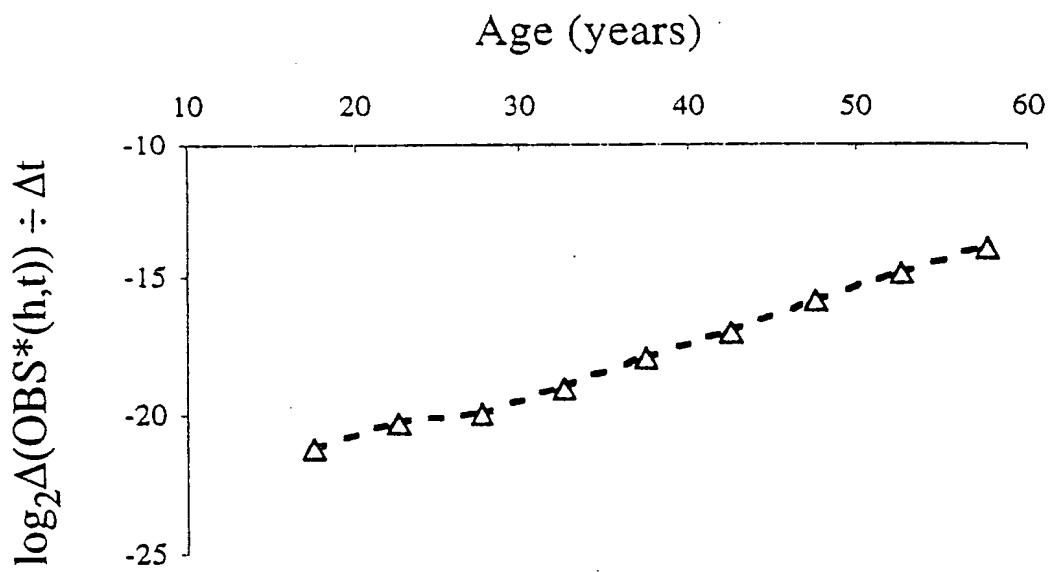


Fig. 17

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Title: Methods of Identifying Point ...
Inventor: William G. Thilly



Fig. 18

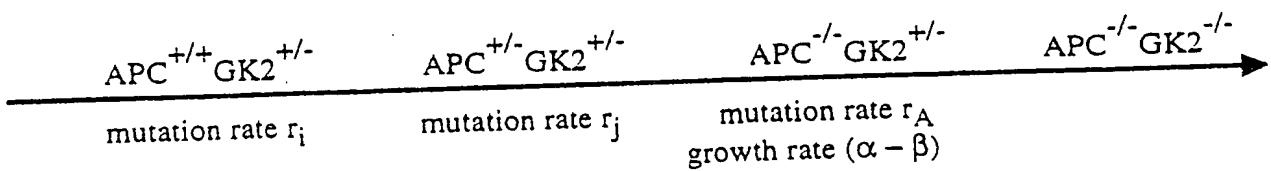




Fig. 19A

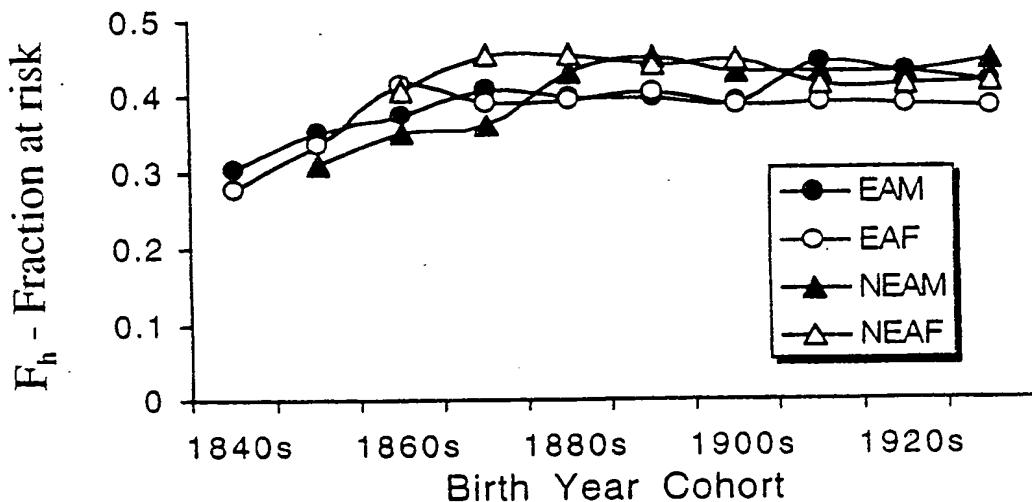


Fig. 19B

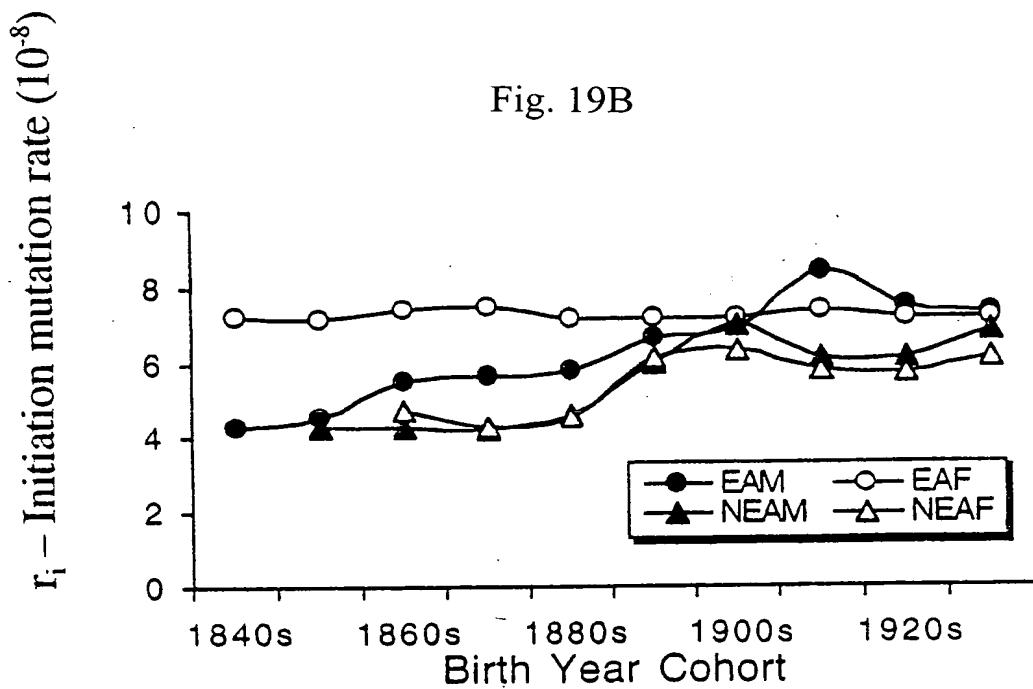




Fig. 20A

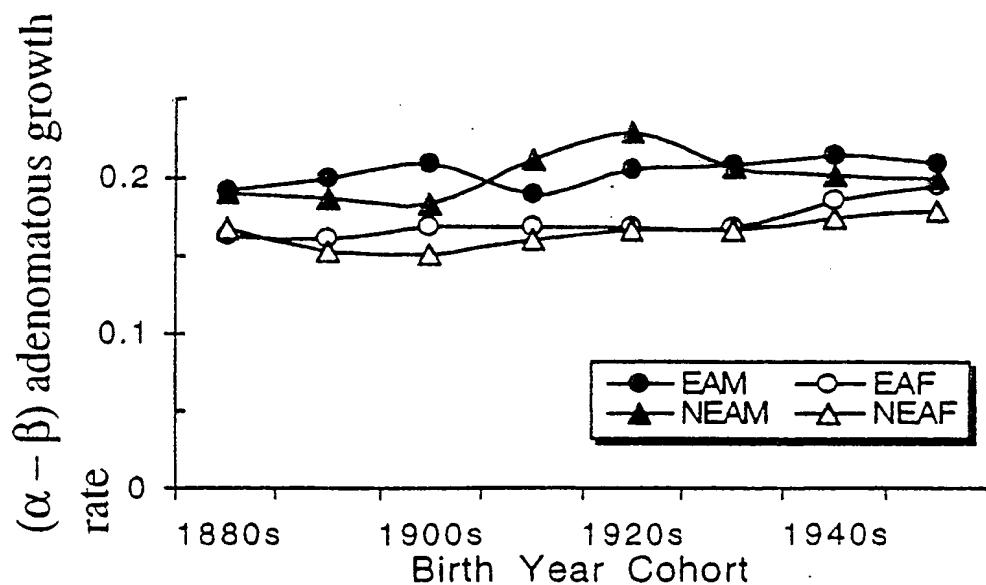
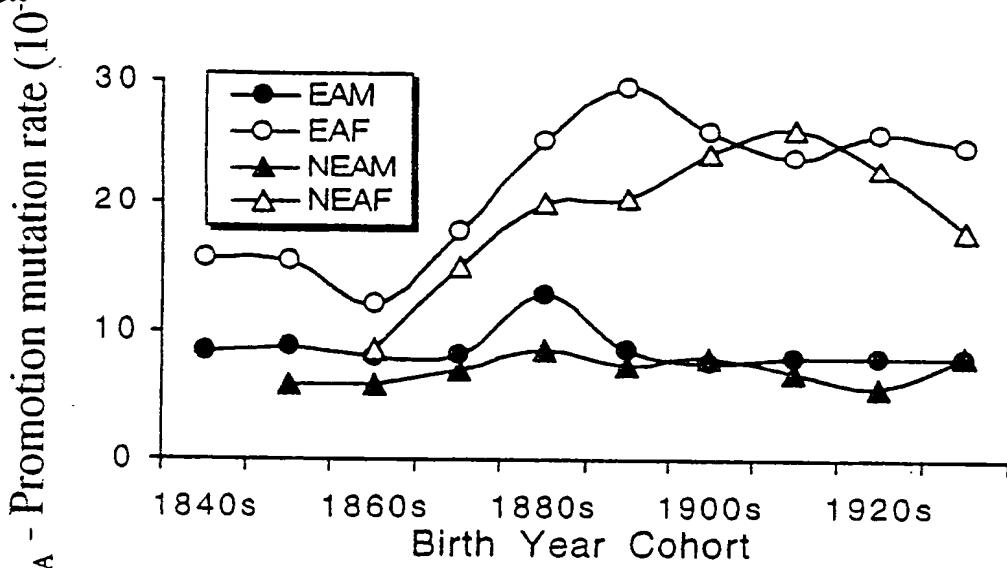


Fig. 20B



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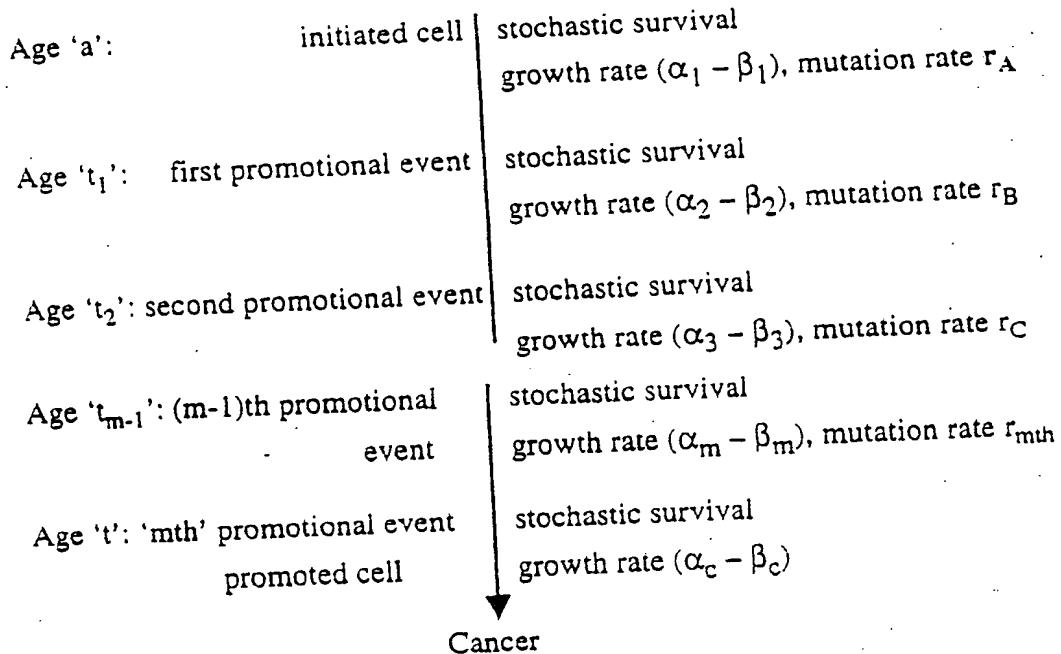


Fig. 21



Fig. 22A

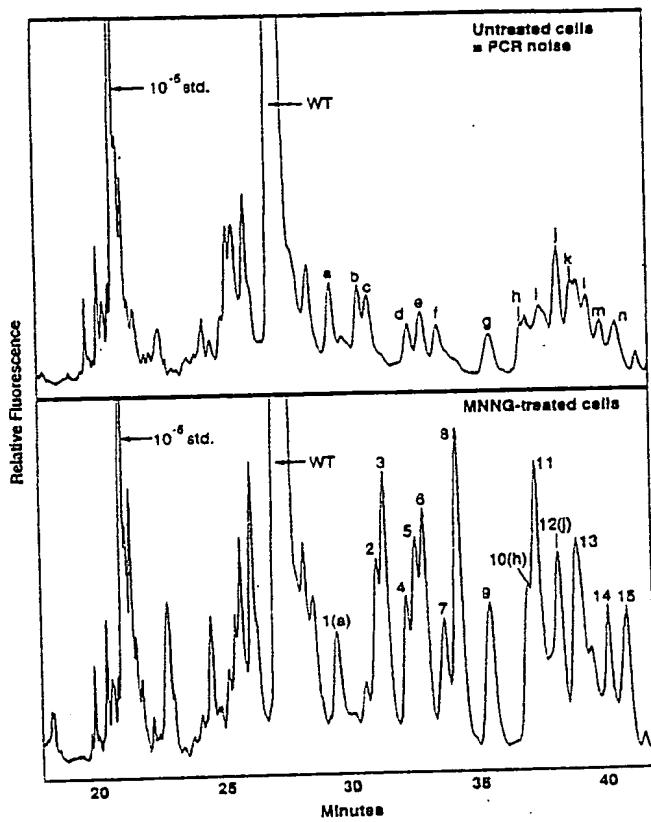
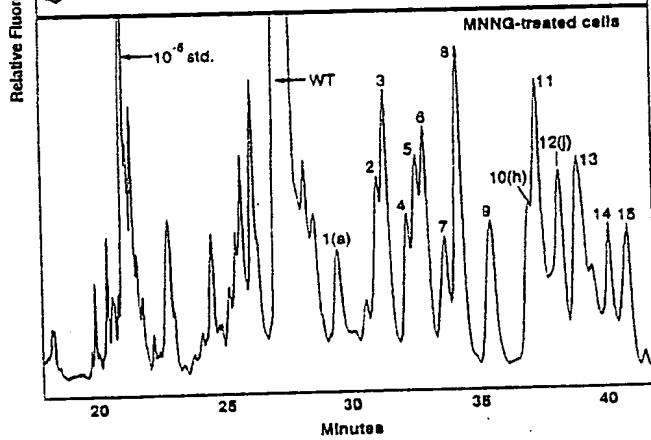


Fig. 22B



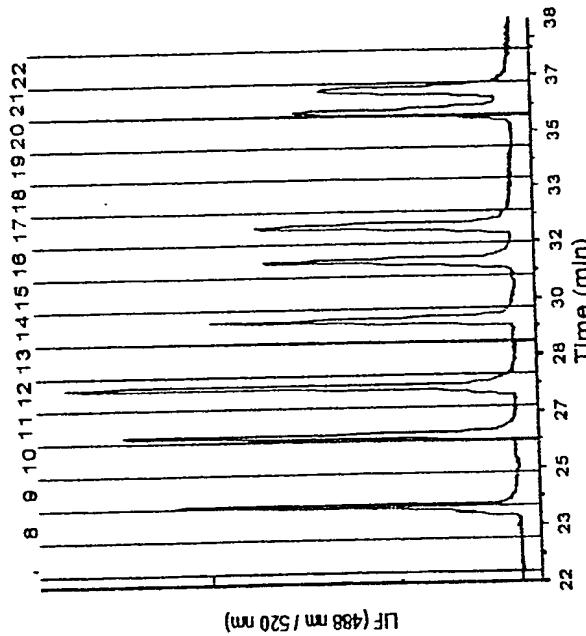


Fig. 23A

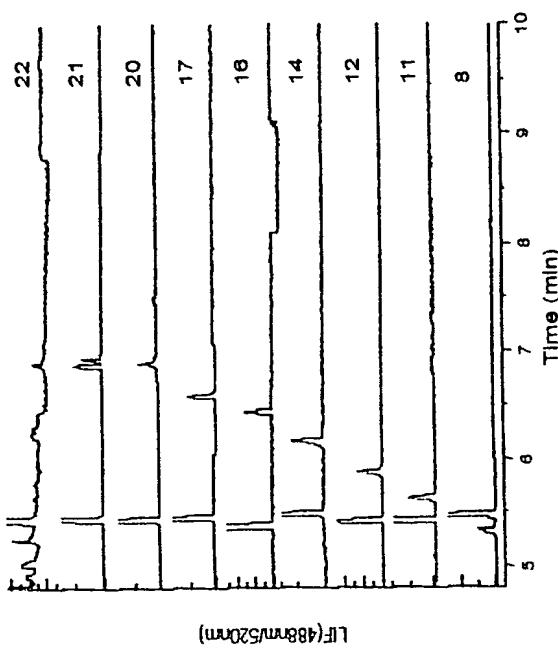


Fig. 23B

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Fig. 24

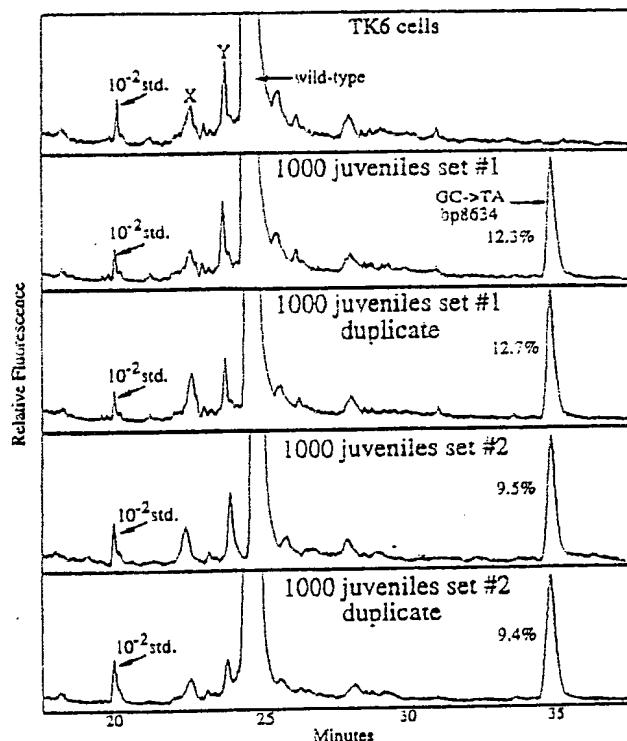




Fig. 25

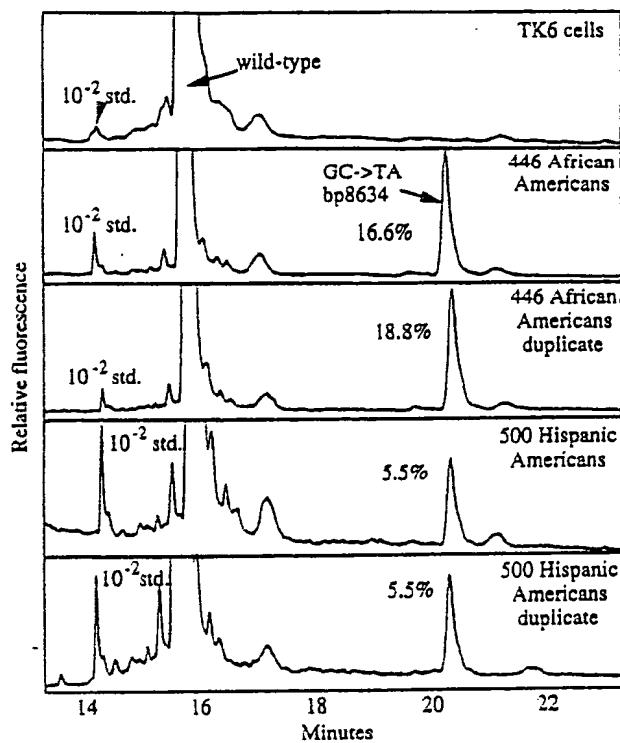


Fig. 26A

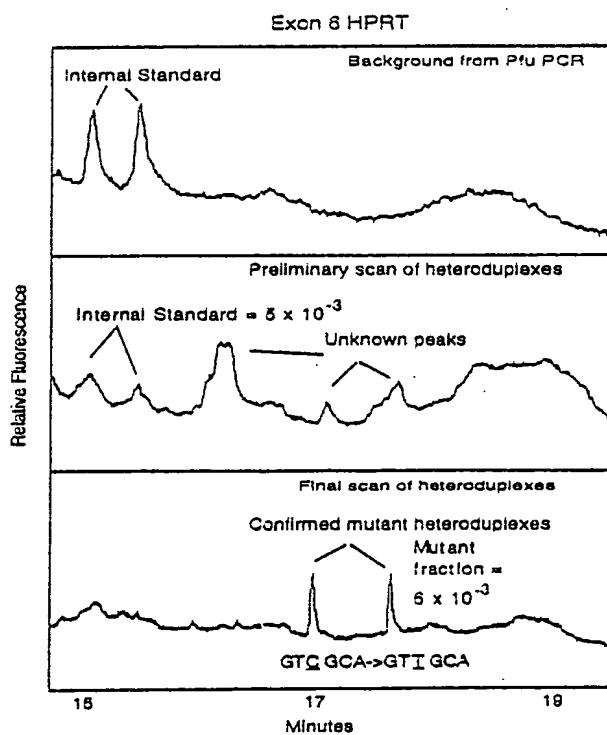




Fig. 27

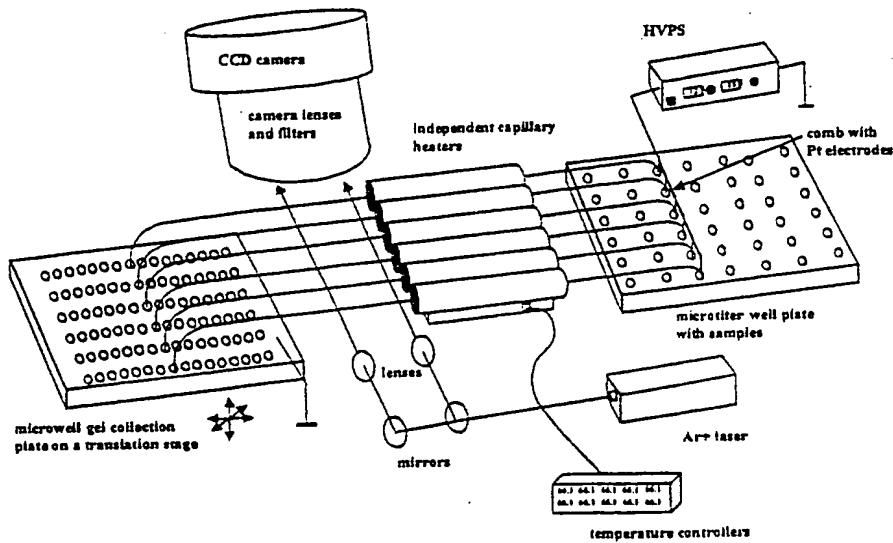
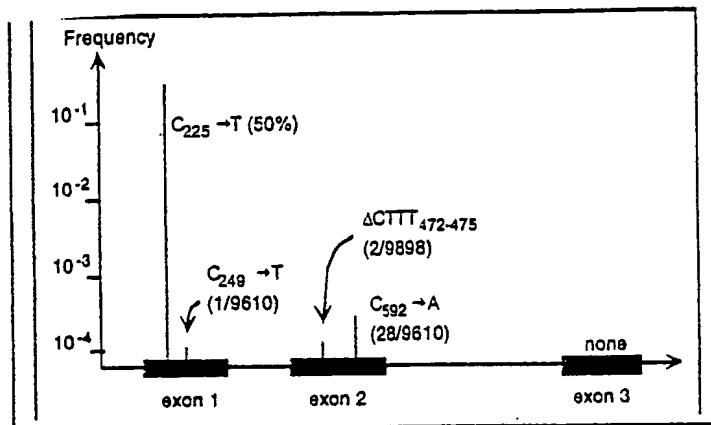


Fig. 28



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Fig. 29

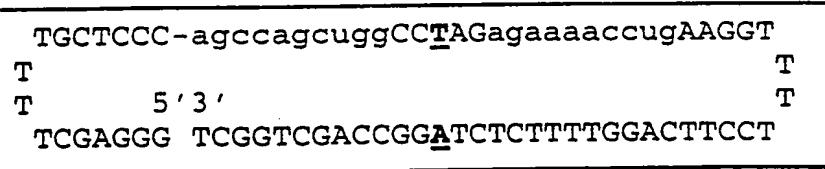
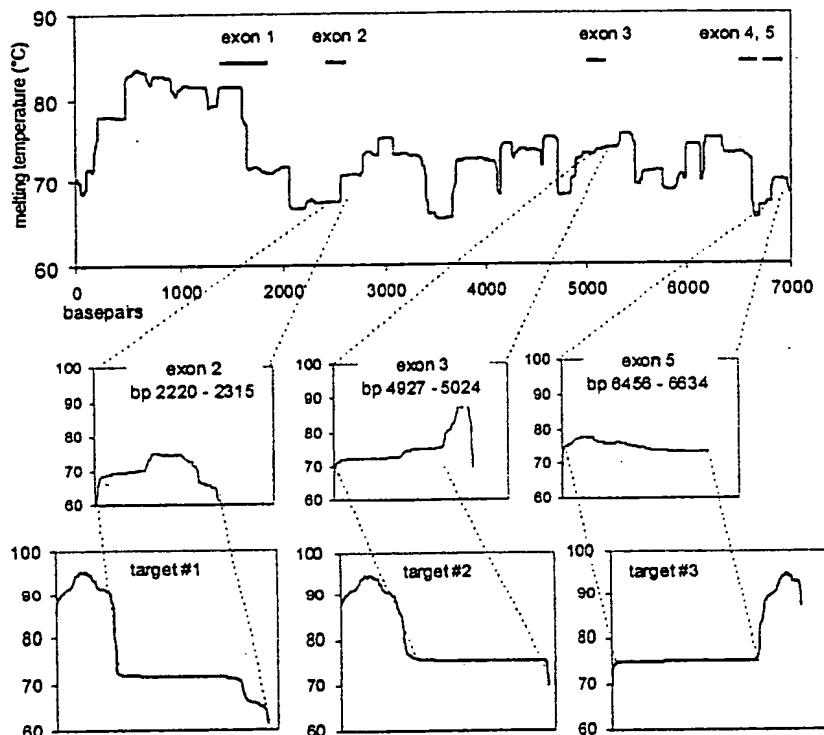


Fig. 30